

CRF Errors Corrected by the STIC Systems Branch
Serial Number: 10/053,530 CRF Processing Date: 04/04/2002
Entered by: J.A. Edited by: J.A. Verified by: J.A. (STIC staff)

OJPE
2/26/2002

Changed a file from non-ASCII to ASCII.

Changed the margins in cases where the sequence text was "wrapped" down to the next line.

Edited a format error in the Current Application Data section, specifically:

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____.

Added the mandatory heading and subheadings for "Current Application Data".

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included:

Deleted extra, invalid, headings used by an applicant, specifically:

Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file;
 page numbers throughout text; other invalid text, such as _____.

Inserted mandatory headings, specifically:

Corrected an obvious error in the response, specifically:

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically:

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

Other: seq 3 - inserted hard return

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

~~HED~~

OIPE

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/10/053,530

DATE: 02/26/2002
 TIME: 12:23:09

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 Output Set: N:\CRF3\02262002\J053530.raw

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2      Hayden-Ledbetter, Martha
3 <120> TITLE OF INVENTION: Binding Domain-Immunoglobulin Fusion Proteins
4 <130> FILE REFERENCE: 390069.401
5 <140> CURRENT APPLICATION NUMBER: US/10/053,530
6 <141> CURRENT FILING DATE: 2002-01-17
7 <150> PRIOR APPLICATION NUMBER: US 09/765,208
8 <151> PRIOR FILING DATE: 2001-01-17
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20 <221> NAME/KEY: V_region
21 <222> LOCATION: (79)..(396)
22 <223> OTHER INFORMATION: light chain variable region for anti-CD20 scFv
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RAW SEQUENCE LISTING
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RAW SEQUENCE LISTING
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96 <223> OTHER INFORMATION: HINGE CYSTEINES (826-829; 844-847; 853-856) MUTATED TO
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 98 PROLINE TO SERINE MUTATION (880-883) IN CH2 DISRUPTS EFFECTOR FUN
 99 CTION
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 116 ccaaaaccca aggacaccctt catgtatctcc cggaccctg aggtcacaatg cgtggcgg 960
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 140 853-856)
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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/053,530

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Input Set : N:\Crf3\02132002\J053530.raw
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STATISTICS SUMMARY
PATENT APPLICATION: US/10/053,530

DATE: 02/26/2002
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Input Set : N:\Crf3\02132002\J053530.raw
Output Set: N:\CRF3\02262002\J053530.raw

Application Serial Number: US/10/053,530

Alpha or Numeric: Numeric

Application Class:

Application File Date: 01-17-2002

Art Unit: OIPE

Software Application: PatentIN3.0

Total Number of Sequences: 38

Total Nucleotides: 14732

Total Amino Acids: 4719

Number of Errors: 0

Number of Warnings: 0

Number of Corrections: 0

MESSAGE SUMMARY